OIPE

 RAW SEQUENCE LISTING
 DATE: 06/19/2001

 PATENI APPLICATION: US/09/755,016
 TIME: 12:22:03

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\06192001\1755016.raw

ENTERED

```
4 <110> APPLICANT: Walke, D. Wade
              Wilganowski, Nathaniel L.
              Donoho, Gregory
      b
              Turner, C. Alexander Jr.
      9 <120 - TIFLE OF INVENTION: Novel Human Proteases and Polynucleotides Encoding the Same
     11 - 150 - Fine REFERENCE: LEX-0114-USA
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/755,016
C--> 13 <141> CURRENT FILING DATE: 2001-06-01
     13 <150> PRIOR APPLICATION NUMBER: US 60/174,686
     14 -(151): PRIOR FILING DATE: 2000-01-06
     16 -:1600- NUMBER OF SEQ ID NOS: 7
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     21 <211> LENGTH: 654
     22 K2125 TYPE: DNA
     23 <213 > ORGANISM: Homo sapien
     25 <400 > SEQUENCE: 1
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                                                                                 b()
     27 ataaaaatta aagcaatcat tattcatcca aacttcattt tqqaatctta tqtaaatgat
                                                                                120
     28 attgeacttt tteacttaaa aaaaqeagtg aggtataatg actatattea geetatttge
                                                                                180
     29 ctaccttttg atgitticca aatcetggae ggaaacacaa agigtittat aagiggeigg
                                                                                240
                                                                                300
     30 qgaaqaacaa aaqaaqaaqq taacqctaca aatattttac aaqatqcaqa aqtqcattat
                                                                                 360
     31 atttetegag agatgtgtaa ttetgagagg agttatgggg gaataaftee taaeaettea
                                                                                420
     eta 2 littigigoag gigalgaaga tygagottiti qalacifgoa qqqqiqacaq igqqqqacca
     33 ttaatgtget aettaeeaga atataaaaga tttt/tgtaa tgggaattae eagttaegga
                                                                                480
     34 catggetgig gtegaagagg titteetiggt gretatatig ggeeateeti etaecaaaag
                                                                                540
     35 tqqotqabaq aqbatttott boat,qbaaqo abtoaaqqoa tabttabtat aaatatttta
                                                                                (x)+41
                                                                                654
     36 cqtqqccaqa foctscatago fittatqffff qfcaloftac tabouaccado atala.
     38 -210> SEQ ID NO: 2
     3 + + 21 ' + LENGTH: 217
     4^{\prime\prime} > 242 \times 1YPE: PRI
     41 - zi3 - oRGANISM Homo sapien
     43 + 400 > SEQUENCE: 2
     44 Met Trp Thr Ala Val Ile Gly Thr Ash Ash Ile His Gly Arg Tyr Pro
     45 1 5
                                    1 ()
     46 His Thr Lys Lys Ile Lys Ile Lys Ala Ile Ile Ile His Prò Asn Phe
                                        z^{(-1)}
                    20
     48 The Lou Glu Ser Iv: Val Asse Ast the Alartee 11 William Co. 1
```

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Input Set : A:\Seqlist.txt

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58 Gly Gly Ile Ile Pro Asn Thr Ser Phe Cys Ala Gly Asp Glu Asp Gly
59
           115
                                120
                                                     125
60 Ala Phe Asp Thr Cys Arg Gly Asp Ser Gly Gly Pro Leu Met Cys Tyr
                           135
62 Leu Pro Glu Tyr Lys Arg Phe Phe Val Met Gly Ile Thr Ser Tyr Gly
                                            155
                       150
64 His Gly Cys Gly Arg Arg Gly Phe Pro Gly Val Tyr Ile Gly Pro Ser
                                        170
                   165
66 Phe Tyr Gln Lys Trp Leu Ihr Glu His Phe Phe His Ala Ser Thr G!n
6.7
                                   185
               180
68 Gly Ilo Leu Thr lle Asn Ile Leu Arg Gly Gln Ile Leu Ile Ala Leu
           195
                                200
70 Cys Phe Val Ile Leu Leu Ala Thr Thr
71
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                           215
73 <210 > SEQ 1D NO: 3
74 - 211 > LENGTH: 1047
75 <212> TYPE: DNA
76 <213 DRGANISM: Homo sapien
78 <400> SEQUENCE: 3
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                                                                             60
                                                                            120
80 gaccaetact egocolotyg aagydacagg otoggoddot dyddygaaco gynggotagt
                                                                            180
81 teccaycayg ctyaygecyt cegcaaqayg ctecggegge ggagggaggg aggggegeat
82 qcaaaqqatt qtqqaacaqc accqc+taaq qatqtqttqc aaqqqtctcq qattatagqq
                                                                            240
                                                                            300
83 ggcaccgaag cacaagctgg cgcatggccg tgggtggtga gcctgcagat taaatatggc
                                                                            360
84 cytqttettg tteatgtatg tgggggaaec etaqtgagag agaqqtqqqt ceteacaqet
                                                                            420
85 geccaetgea etaaagaere tagegateet ttaatgtgga eagetgtgat tggaactaat
86 aatatacatq gacqctatcc tcataccaag aagataaaaa ttaaagcaat cattattcat
                                                                            480
                                                                            540
87 ccaaacttea ttttggaate ttatgtaaat gatattgeae ttttteaett aaaaaaagea
88 qtgaqqtata atqactatat tcaqcctaft fqcctacctt ttqatqtftt ccaaatcctq
                                                                            600
                                                                            660
89 gacqqaaaca caaaqtqttt tataaqtqqc tqqqqaaqaa caaaaqaaqa aqqtaacqct
                                                                            720
90 acaaatattt tacaaqatqc aqaaqtqcat tatafttctc qaqaqatqtq taatfctqaq
                                                                            780
91 aqqaqttatq qqqqaataat tootaacact toattttqtq caqqtqatqa aqatqqaqct
                                                                           840
92 fitgatacti gcaggggtga cagtggggga ccaltaatgi gctacttacc agaatataaa
                                                                           41111
9s agailtiitty taatyggaat taccayltac yyacatyyct ylygicgaay agailticci
94 gatatetata tigagoreato ottobaccaa aagtagootga cagagoatit obtocalaca
                                                                           198512
                                                                           . 2
The agractically gratactiae latalatati titaegisgen agaiteeteat agentitatut
🤲 titqteafet täctageaac aacataa
                                                                           1047
^{68} + 210; SEQ ID NO: 4
99 + 211> LENGTH: 348
100 <212> TYPE: PRT
101 - 213 - ORGANISM Homo sapien
103 - 400 - SEQUENCE: 4
104 Met Arq Leu Gly Leu Leu Ser Val Mila to Liber 13. The transfer
```

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 PATENT APPLICATION
 US/09/755,016
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111 50
                      5.5
112 Gly Thr Ala Pro Lea Lys Asp Val Leu Gln Gly Ser Arg Ile Ile Gly
         70
114 Gly Thr Glu Ala Gln Ala Gly Ala Trp Pro Irp Val Val Ser Leu Gln
          8.5
                                90
lle lie Lys Tyr Gly Arg Val Leu Val His Val Cys Gly Gly Thr Leu Val
117 100
                          105
118 Arg Giu Arg Irp Val Leu Thr Ala Ala His Cys Thr Lys Asp Ala Ser
11. 115 120 125
120 Asp Pro Leu Met Trp Thr Ala Val Ile Gly Thr Asn Asn Ile His Cly
121 130 135
                             140
122 Arg Tyr Pro His Thr Lys Lys lle Lys Ile Lys Ala Ile Ile Ile His
12 + 145
                   150
                                   155
124 Pro Asn Phe Ile Leu Glu Ser Tyr Val Asn Asp Ile Ala Leu Phe His
125
                165
                                170
126 Leu Lys Lys Ala Val Arg Tyr Asn Asp Tyr Ile Gln Pro Ile Cys Leu
127 180
                            185
128 Pro Phe Asp Val Phe Gln Ile Leu Asp Gly Asn Thr Lys Cys Phe Ile
129 195
                         200
130 Ser Gly Trp Gly Arg Thr Lys Glu Glu Gly Asn Ala Thr Asn Ile Leu
131 210
                     215
                                      220
132 Gln Asp Ala Glu Val His Tyr Ile Ser Arg Glu Met Cys Asn Ser Glu
                  230
                                   235
134 Arg Ser Tyr Gly Gly Ile Ile Pro Asn Thr Ser Phe Cys Ala Gly Asp
                245
                                250
136 Glu Asp Gly Ala Phe Asp Thr Cys Arg Gly Asp Ser Gly Gly Pro Leu
137 260
                            265
                                             270
138 Met Cys Tyr Leu Pro Glu Tyr Lys Arg Phe Phe Val Met Gly Ile Thr
139 275
                         280
140 Ser Tyr Gly His Gly Cys Gly Arg Arg Gly Phe Pro Gly Val Tyr Ile
141 290
                      295
142 Gly Pro Ser Phe Tyr Gln Lys II; Ieu Ihr Glu His Phé Phe His Ala
                   310
                                   $ 1 1)
144 Ser thr Gln Gly He Ieu Thr Ile Ash Ile Ieu Arg Gly Gln He Ieu
145
                     ₹⊀⊓
146 He Ala Leu Cys Phe Val The Leu Leu Ala Thr The
147 340
149 <210> SEQ 1D NO: 5
150 < 211 > LENGIH 867
151 <212> TYPE: DNA
152 - 213 > ORGANISM: Homo sapien
154 - 4000 SEQUENCE: 5
155 atgragaatt gtogrageade accondition ook of statistic vicinities ook o
```

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RAW SEQUENCE LISTINGPATENT APPLICATION: US/09/755,016

DATE: 06/19/2001
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Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\06192001\1755016.raw

162	gace	ggaaa	аса (caaac	qtgti	tt. ta	at.aa	qtgg	r tg	qqqaa	адаа	caaa	aagaa	aga a	aggtá	aacget.
163	acad	aatai	t t.t.	tacaa	agato	go ao	gaagt	tgcal	t. t.ar	tatt	tata	gaga	agato	gtg	t.aat, t	ctgag
1 to 4	agga	agtta	at.g -	gaaga	aataa	at to	cctaa	acact	t. t.c.	ait, titi ti	Lgtg	cago	gtiga	tga a	agato	ggaget
165	t.t.t.	jata	itt.	geage	gggto	ja ca	agtgo	gggg	a cc	att.aa	at.qt	get.a	act.t.a	acc a	agaat	ataaa
156	agat	t t.t.t	ttg :	taato	qqqaa	at ta	acca	gt tac	2 99	acato	gget	gtg	gtoga	ववप्र	aggtit	ttcct
167	ggt	at.cta	ata :	ttgg	gedat	tic ict	tteta	accaa	a da	gt.qq	etga	caga	agidal	ttt d	ette	catgca
168	àqca	actca	aaq	gcata	actta	ac ta	ataaa	atati	t t.t.	acgto	19cc	agat	taat	dat a	agett	tatgt
169 tiligicatet taetageaac aacataa																
171 <210> SEQ ID NO: 6																
172 <211> LENGTH: 288																
173 -CE12 - TYPE: PRI																
	74 ×213> ORGANISM: Homo sapien 76 ×400> SEQUENCE: 6															
						on l	. 1								.2.1.	a
		GIN	Asn	Cys	GLY	Thr	Ala	Pro	Leu	Lys 10	Asp	vai	Leu	GIN	61y 15	ser
178	1	T1 -	Tlo	Gly	.,	The	Clo	A Lo	Ola		Clir	A 1 a	Trin	Dro		V > 1
$\frac{1}{180}$	Arg	* T.G.	rre	-50 -513	этА	1111	GLU	ALd	25	Ald	GLY	Ald	пр	30	пр	val
	Val	Cor	Lou	Gin	Tlo	LVC	Tur	Clu		Val	Lau	Wal	иiс		Cura	Clv
182	vai	ser	35	9111	TTG	пуз	1 7 1	40	Arg	val	Ltsu	Val	45	vai	'- Y 3	GIY
	Glv	Thr		Val	Ara	Glu	Ara		Va l	Len	Thr	Ala	-	His	(779	Thr
184	·) - 1	50	LINEGE	• • • •	9	010	55	116	, 41	БСС		60	11.1 0		~ I W	
	Lys		Ala	ser	Asp	Pro		Met.	anT	Thr	Ala		Ile	Gly	Thr	Asn
186	1	1				70					7.5			1		80
187	Asn	He	His	Gly	Arg	Tyr	Pro	His	Thr	Lys	Lys	Пe	Lys	He	Lys	Ala
188					85					90					95	
189	11e	116	He	His	Pro	Asn	Phe	Ile	Leu	Glu	ser	Tyr	Val	Asn	Asp	Ile
190				100					105					110		
191	Ala	Leu	Phe	His	Leu	Lys	Lys	Ala	Val	Arg	Tyr	Asn	Asp	туг	Ile	Gln
192			115					120					125			
	Pro		Cys	Leu	Pro	Phe		Val	Phe	Gln	I, f.G		Asp	Gly	Asn	Thr
194		130				. 5	135					140				
		Cys	Pho	He	198		144	GLY	ताप	Thir		Glu	Glu	GTY	Asn	
	.45				* 1	150					155				• 1	166
	1111	ASII	110	Len		ASP	Ala	(, 11	Val		1 !	1110	201	75 I U		[-]+ ,]
198				Giu	165		4		(11	1 1	1 0	Dia .	7	: 1. s	175	111.5
201	E 7 25	nsh	26.1	180	AIG	201	1 > 1	CF L y	185	+ 1 + 1	1 +,	1.1.1	25.25.11	190	124.1	11.0
	(11.0	7. L.a	C1 · ·	Asp	(*10	Λαι	C1	Δla		Λατι	The	1	λεσ		Aen	Sar
201	. 12	MIG	195	ush,	OIU	ust.	017	200	1 116	nol,	1111	15	205	OIY	nop	101
	Glv	Glu		Leu	Met	Cvs	Evr		Pro	Glu	Tire	Lvs		Pho	Pho	Val
204	y	210	1 1 ()	131 (1	A LN Text	· / · ·	215	4.1 14	/		* . *	1.72()	,			
	Mest		H	Thi	Ser	Tur	_	His	$G \Vdash \nabla$	CVS	(; [·:		Ara	G1v	Phe	Pro
	225	* * 1				1 3 6 1	,,,		* * 7	7.5		,	,			

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/755,016

DATE 06/19/2001 TIME: 12:22:03

1200

1260

1286

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\06192001\I755016.raw

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.:21	aaggggctgg	ctogeogeca	tettgeteac	cagedtecaa	aatgcggctg	gggctdctga	120							
222	gegtggeget	gt.tgt/tgtg	qqqagctctc	acttatactc	agaccactac	tagacatatg	180							
223	qaagqcacag	gataygadaa	t.cqccggaac	cyqcygctay	ttcccaqcag	getgaggéeg	240							
124	teegeaagag	geteeggega	cddagddadd	gaggggga	t.gloaaaggat	tgtggäācay	300							
225	caecgettaa	ggatgtgttg	caagggtete	ggattatagg	gggcaccgaa	gdadaagdtg	360							
226	gegeatggee	gtggqtggtg	agcotgoaga	ttaaatatgq	cogtyttett	gttcatgtat	4.20							
227	gtgggggaac	cctagtgaga	gagaggtggg	tectcacage	tgcccactgc	actaaagacg	4 H O							
228	ctagegatee	tttaatgtgg	acagetqtga	ttggaactaa	taatatacat	ggaegetate	540							
229	ctcataccaa	gaagataaaa	attaaaycaa	tcattattca	tecaaacttc	attttggaat	() () i)							
230	cttatgtaaa	tgatattgca	ctttttcact	taaaaaaagc	agtgaggtat	aatyactata	f) F. ()							
231	ttdagddtat	ttgcctacct	tttgatgttt	tecaaateet	ggacggaaac	acaaaqtgtt	720							
232	ttataagtgg	ct.ggygaaga	acaaaagaag	aaggtaacgc	tacaaatatt	ttacaagatg	780							
233	cagaagtgca	ttatatttct	cgagagatgt	gtaattetga	gaggagttat	gggggaataa	840							
234	ttectaacac	ttcaitttgt	qcaggtqatg	aagatggagc	tittgatact	tgcaggggtg	(1(11)							
235	acagtggggg	ancattaatg	tgctacttac	cagaatataa	aagatttitt	gtaatgggaa	(44.1)							
236	ttaccagtta	cggacatggc	tgtggtcgaa	gaggttttcc	tggtgtctat	attgggccat	1020							
237	ccttctacca	aaagtggctg	acagagcatt	tettecatge	aagcactcaa	ggcatactta	1080							
238	ctataaatat	tttacgtggc	cagatectea	tagctttatg	t*ttgtcatc	ttactagcaa	1140							

239 caacataaay aaattotgaa ggotttoata tottfatttt geattgtgto ootttotatg

240 tictatataa igaacatcai tialiictict agcaattaai igcclacati agagattica

241 tgtgaacatt ttatgggcta taaata

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/755,016

DATE: 06/19/2001 TIME: 12:22:04

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\06192001\1755016.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:i3 M:271 C: Current Filing Date differs, Replaced Current Filing Date